Diversity and abundance of bacteria
in surface seawater of the Changjiang Estuary and its adjacent areas

PHYLOGENETIC TREES

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Fig. S1. Phylogenetic tree of Proteobacteria according to 16S rDNA gene sequences. Alpha Proteobacterium, Beta Proteobacterium, Gamma Proteobacterium, Delta Proteobacterium and Epsilon Proteobacterium are α, β, γ, δ and ε, respectively (in different colors).
Phylogenetic tree of other bacteria according to 16S rDNA gene sequences.